

Resource Summary Report

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Antibodypedia

RRID:SCR_012782

Type: Tool

Proper Citation

Antibodypedia (RRID:SCR_012782)

Resource Information

URL: <http://www.antibodypedia.com/>

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Description: Open-access database of antibodies against human proteins developed through collaboration between Antibodypedia AB and the Nature Publishing Group. It aims to provide the scientific community and antibody distributors alike with information on the effectiveness of specific antibodies in specific applications--to help scientists select the right antibody for the right application. Antibodypedia's mission is to promote the functional understanding of the human proteome and expedite analysis of potential biomarkers discovered through clinical efforts. To this end, they have developed an open-access, curated, searchable database containing annotated and scored affinity reagents to aid users in selecting antibodies tailored to specific biological and biomedical assays. They envisage Antibodypedia as a virtual repository of validated antibodies against all human, and ultimately most model-organism, proteins. Such a tool will be exploitable to identify affinity reagents to document protein expression patterns in normal and pathological states and to purify proteins alone and in complex for structural and functional analyses. They hope to promote characterization of the roles and interplay of proteins and complexes in human health and disease. They encourage commercial providers to submit information regarding their inventory of antibodies with links to quality control data. Independent users can submit their own application-specific experimental data using standard validation criteria (supportive or non-supportive) developed with the assistance of an international advisory board recruited from academic research institutions. Users can also comment on specific antibodies without submitting validation data.

Synonyms: Antibodypedia / Nature

Resource Type: data or information resource, database

Defining Citation: [PMID:18667413](#), [PMID:18767878](#)

Keywords: cell biology, antibody, protein, human, reagent, model organism, non-human primate, FASEB list

Funding: Antibodypedia AB ;
Nature Publishing Group ;
European Union 6th framework - ProteomeBinders ;
Human Antibody Initiative ;
HUPO - Human Proteome Organisation

Availability: The community can contribute to this resource

Resource Name: Antibodypedia

Resource ID: SCR_012782

Alternate IDs: nif-0000-22918, OMICS_01770

Record Creation Time: 20220129T080312+0000

Record Last Update: 20260905T133203+0000

Ratings and Alerts

No rating or validation information has been found for Antibodypedia.

No alerts have been found for Antibodypedia.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 45 mentions in open access literature.

Listed below are recent publications. The full list is available at [PRECISE-TBI](#) .

Robinson JP, et al. (2026) Flow Cytometry: Advances, Challenges and Trends. BioEssays : news and reviews in molecular, cellular and developmental biology, 48(1), e70091.

Kafita D, et al. (2026) Decoding the dark genome reveals its organisation into modular disease networks. Scientific reports, 16(1).

Liu J, et al. (2025) Comprehensive characterization of multi-omics landscapes between gut microbial metabolites and the druggable genome in sepsis. Frontiers in immunology, 16, 1597676.

Broderick K, et al. (2025) Human protein interaction networks of ancestral and variant SARS-CoV-2 in organ-specific cells and bodily fluids. Nature communications, 16(1), 5784.

Bonilla DL, et al. (2024) The Power of Reagent Titration in Flow Cytometry. Cells, 13(20).

- Morey A, et al. (2024) The EV antibody database: An interactive database of curated antibodies for extracellular vesicle and nanoparticle research. *Extracellular vesicle*, 3.
- Kahn RA, et al. (2024) Antibody characterization is critical to enhance reproducibility in biomedical research. *eLife*, 13.
- Geyer PE, et al. (2024) The Circulating Proteome?Technological Developments, Current Challenges, and Future Trends. *Journal of proteome research*, 23(12), 5279.
- Li J, et al. (2024) Single-cell and bulk RNA-sequence identified fibroblasts signature and CD8 + T-cell - fibroblast subtype predicting prognosis and immune therapeutic response of bladder cancer, based on machine learning: bioinformatics multi-omics study. *International journal of surgery (London, England)*, 110(8), 4911.
- Laflamme C, et al. (2021) Opinion: Independent third-party entities as a model for validation of commercial antibodies. *New biotechnology*, 65, 1.
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- Swier VJ, et al. (2020) Validating indicators of CNS disorders in a swine model of neurological disease. *PloS one*, 15(2), e0228222.
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- Proietti Onori M, et al. (2018) The intellectual disability-associated CAMK2G p.Arg292Pro mutation acts as a pathogenic gain-of-function. *Human mutation*, 39(12), 2008.
- Meliopoulos VA, et al. (2018) Although it's painful: The importance of stringent antibody validation. *PLoS pathogens*, 14(1), e1006701.
- Hung YF, et al. (2018) Endosomal TLR3, TLR7, and TLR8 control neuronal morphology through different transcriptional programs. *The Journal of cell biology*, 217(8), 2727.
- Chen Z, et al. (2018) Current applications of antibody microarrays. *Clinical proteomics*, 15, 7.
- Zhang W, et al. (2018) Generation of a monoclonal antibody recognizing the heavily glycosylated CD45 protein and its application on identifying circulating tumor cells. *PloS one*, 13(2), e0192506.